

## SEQUENCE LISTING

<110> GARABEDIAN, Michael  
TANEJA, Samir  
HITTELMAN, Adam  
MARKUS, Steven

<120> METHOD FOR SCREENING TRANSCRIPTIONAL COREGULATORY PROTEINS OF TRANSCRIPTION FACTORS, AND ANDROGEN RECEPTOR TRANSCRIPTIONAL COREGULATORY PROTEINS AS TARGETS FOR ANDROGEN RECEPTOR-DEPENDENT DISEASES

<130> GARABEDIAN=1.1A

<140> NOT YET ASSIGNED

<141> 2001-03-26

<150> 60/225,618

<151> 2000-08-15

<150> 60/191,768

<151> 2000-03-24

<160> 20

<170> PatentIn version 3.0

<210> 1

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<400> 1

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<400> 2

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20           25           30
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Arg Lys Val Leu Asp His Arg Asp Lys Val Tyr Glu Gln Leu Ala Lys
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Ser Glu Leu Tyr Met Gln Val Asp Leu Gly Cys Asn Phe Phe Val Asp		
65	70	75
Thr Val Val Pro Asp Thr Ser Arg Ile Tyr Val Ala Leu Gly Tyr Gly		
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Phe Phe Leu Glu Leu Thr Leu Ala Glu Ala Leu Lys Phe Ile Asp Arg		
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Lys Ser Ser Leu Leu Thr Glu Leu Ser Asn Ser Leu Thr Lys Asp Ser		
	115	120
Met Asn Ile Lys Ala His Ile His Met Leu Leu Glu Gly Leu Arg Glu		
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<212> PRT

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Ala Gly His Asp Ala Pro Arg Arg Thr Arg Pro Ser Ala Arg Lys Pro  
35 40 45

Leu Ser Cys Phe Ser Arg Arg Pro Met Pro Thr Arg Glu Pro Pro Lys  
50 55 60

Thr Arg Gly Ser Arg Gly His Leu His Thr His Pro Pro Gly Pro Gly  
65 70 75 80

Pro Pro Leu Gln Gly Leu Ala Pro Arg Gly Leu Lys Thr Ser Ala Pro  
85 90 95

Arg Pro Pro Cys Gln Pro Gln Pro Gly Pro His Lys Ala Lys Thr Lys  
100 105 110

Lys Ile Val Phe Glu Asp Glu Leu Leu Ser Gln Ala Leu Leu Gly Ala  
115 120 125

Lys Lys Pro Ile Gly Ala Ile Pro Lys Gly His Lys Pro Arg Pro His  
130 135 140

Pro Val Pro Asp Tyr Glu Leu Lys Tyr Pro Pro Val Ser Ser Glu Arg  
145 150 155 160

Glu Arg Ser Arg Tyr Val Ala Val Phe Gln Asp Gln Tyr Gly Glu Phe  
165 170 175

Leu Glu Leu Gln His Glu Val Gly Cys Ala Gln Ala Lys Leu Arg Gln  
180 185 190

Leu Glu Ala Leu Leu Ser Ser Leu Pro Pro Pro Gln Ser Gln Lys Glu  
195 200 205

Ala Gln Val Ala Ala Arg Val Trp Arg Glu Phe Glu Met Lys Arg Met  
210 215 220

Asp Pro Gly Phe Leu Asp Lys Gln Ala Arg Cys His Tyr Leu Lys Gly  
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 ggaaggcgcc cggcaaggag gcggacaagc ggagcaggcc aacgagacgc gcgcacccac 300  
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 <212> PRT  
 <213> Human

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 50 55 60  
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 65 70 75 80  
 Glu Thr Lys Val Leu Gly Ala Leu Leu Phe Val Lys Gly Ala Val Trp  
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 Lys Ala Leu Phe Gly Lys Glu Ala Asp Lys Leu Glu Gln Ala Asn Asp  
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 Asp Ala Arg Thr Phe Tyr Ile Ile Glu Arg Glu Pro Leu Ile Asn Thr  
 115 120 125  
 Tyr Ile Ser Val Pro Lys Glu Asn Ser Thr Leu Asn Cys Ala Ser Phe  
 130 135 140

Thr Ala Gly Ile Val Glu Ala Val Leu Thr His Ser Gly Phe Pro Ala  
145 150 155 160

Lys Val Thr Ala His Trp His Lys Gly Thr Thr Leu Met Ile Lys Phe  
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Glu Glu Ala Val Ile Ala Arg Asp Arg Leu Glu Gly Arg  
180 185

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<211> 126  
<212> DNA  
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caagga 126

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taaaggacga acctgatctc ttatactagt atccttaatc attttttattg ccacaactaa 240  
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<213> Human
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<400> 10
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Arg Ser Thr Asp His Ser Ile Ser Pro Ser Ile Asp Pro His Leu Gln
          20          25          30
Ile Ser His Gln Gln Pro Thr Asn His His Pro Thr Met Thr Asn Gln
          35          40          45
Thr Asn Leu Lys Thr Asn Asp Asn His Thr Gln His
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<210> 11
<211> 1918
<212> DNA
<213> Human
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<211> 252
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<213> Human

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<400> 12
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          20           25           30

Gly Arg Ala Phe Ser Asp Arg Ser Ser Leu Thr Phe His Gln Ala Ile
          35           40           45

His Thr Gly Glu Lys Pro Tyr Lys Cys His Glu Cys Gly Lys Val Phe
          50           55           60

Arg His Asn Ser Tyr Leu Ala Thr His Arg Arg Ile His Thr Gly Glu
65           70           75           80

Lys Pro Tyr Lys Cys Asn Glu Cys Gly Lys Ala Phe Ser Met His Ser
          85           90           95

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Asn Leu Thr Thr His Lys Val Ile His Thr Gly Glu Lys Pro Tyr Lys  
 100 105 110  
 Cys Asn Gln Cys Gly Lys Val Phe Thr Gln Asn Ser His Leu Ala Asn  
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 His Gln Arg Thr His Thr Gly Glu Lys Pro Tyr Arg Cys Asn Glu Cys  
 130 135 140  
 Gly Lys Ala Phe Ser Val Arg Ser Ser Leu Thr Thr His Gln Ala Ile  
 145 150 155 160  
 His Thr Gly Lys Lys Pro Tyr Lys Cys Asn Glu Cys Gly Lys Val Phe  
 165 170 175  
 Thr Gln Asn Ala His Leu Ala Asn His Arg Arg Ile His Thr Gly Glu  
 180 185 190  
 Lys Pro Tyr Arg Cys Thr Glu Cys Gly Lys Ala Phe Arg Val Arg Ser  
 195 200 205  
 Ser Leu Thr Thr His Met Ala Ile His Thr Gly Glu Lys Arg Tyr Lys  
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 <211> 8588  
 <212> DNA  
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Cys Ala Val Cys Asn Lys Phe Thr Thr Asp Asn Leu Asp Met Leu Gly
          35          40          45
Leu His Met Asn Val Glu Arg Ser Leu Ser Glu Asp Glu Trp Lys Ala
          50          55          60
Val Met Gly Asp Ser Tyr Gln Cys Lys Leu Cys Arg Tyr Asn Thr Gln
          65          70          75          80
Leu Lys Ala Asn Phe Gln Leu His Cys Lys Thr Asp Lys His Val Gln
          85          90          95
Lys Tyr Gln Leu Val Ala His Ile Lys Glu Gly Gly Lys Ala Asn Glu
          100          105          110
Trp Arg Leu Lys Cys Val Ala Ile Gly Asn Pro Val His Leu Lys Cys
          115          120          125
Asn Ala Cys Asp Tyr Tyr Thr Asn Ser Leu Glu Lys Leu Arg Leu His
          130          135          140
Thr Val Asn Ser Arg His Glu Ala Ser Leu Lys Leu Tyr Lys His Leu
          145          150          155          160
Gln Gln His Glu Ser Gly Val Glu Gly Glu Ser Cys Tyr Tyr His Cys
          165          170          175
Val Leu Cys Asn Tyr Ser Thr Lys Ala Lys Leu Asn Leu Ile Gln His
          180          185          190
Val Arg Ser Met Lys His Gln Arg Ser Glu Ser Leu Arg Lys Leu Gln
          195          200          205
Arg Leu Gln Lys Gly Leu Pro Glu Glu Asp Glu Asp Leu Gly Gln Ile
          210          215          220
Phe Thr Ile Arg Arg Cys Pro Ser Thr Asp Pro Glu Glu Ala Ile Glu
          225          230          235          240

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Thr Asp Ser Pro Ala Thr Ser Lys Arg Ile Ser Phe Pro Gly Ser Ser
      275                      280                      285

Glu Ser Pro Leu Ser Ser Lys Arg Pro Lys Thr Ala Glu Glu Ile Lys
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Pro Glu Gln Met Tyr Gln Cys Pro Tyr Cys Lys Tyr Ser Asn Ala Asp
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Val Asn Arg Leu Arg Val His Ala Met Thr Gln His Ser Val Gln Pro
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Met Leu Arg Cys Pro Leu Cys Gln Asp Met Leu Asn Asn Lys Ile His
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Glu Lys Leu Ile Met Thr Val Thr Thr Pro Glu Met Val Met Pro Ser
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Asn Ile Leu Pro Ser Ala Ser Thr Glu Gln Ser Gly Asp Leu Lys Pro
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Ser Pro Ala Asp Pro Gly Ser Val Arg Glu Asp Ser Gly Phe Ile Cys
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Trp Lys Lys Gly Cys Asn Gln Val Phe Lys Thr Ser Ala Ala Leu Gln
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Asp Arg His Val Tyr Lys Tyr Arg Cys Asn Gln Cys Ser Leu Ala Phe
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Lys Thr Ile Glu Lys Leu Gln Leu His Ser Gln Tyr His Val Ile Arg
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Ala Leu Lys Lys His Leu Glu Thr Ser His Leu Glu Leu Ser Glu Ala
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Asp Ile Gln Gln Leu Tyr Gly Gly Leu Leu Ala Asn Gly Asp Leu Leu
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 Pro Thr Ser Ser Pro Asp Asn Lys Pro Phe Lys Cys Asn Thr Cys Asn  
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 835 840 845  
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 His Phe Pro Met Thr Thr Glu Thr Leu Leu Gln Leu Gln Gln Gln Gln  
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1895						1900					1905			
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1910						1915					1920			
Val	Asn	Thr	Ala	Ile	Thr	Asp	Thr	Thr	Thr	Gly	Asp	Glu	Gly	Asn
1925						1930					1935			
Ala	Asp	Asn	Asp	Ser	Ala	Thr	Gly	Ile	Ala	Thr	Glu	Thr	Lys	Ser
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Asp	Tyr	Ser	Glu	Thr	Ser	Ser	Leu	Ala	Asp	Pro	Cys	Ser	Pro	Ser
2000						2005					2010			
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2015						2020					2025			
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 Gly Gln Ser Val Val Asn Leu Gln Glu Met Val Leu His Val Pro  
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 2585 2590 2595  
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 Cys Glu Ser Ala Leu Cys Gly Glu Glu Ala Leu Ser Gln His Leu  
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 Asn Ala Lys Glu His Pro Ser Leu Leu Pro His Ser Ala Cys Phe  
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 Asn Asp Ser Pro Pro Pro Pro Ser Ala Ala Ala Pro Ser Ser Ala  
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 2690 2695 2700  
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